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Rec'd PCT/PTO 09 NOV 2001

SEQUENCE LISTING

<110> TOPOROIK, Amir et al.

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<140> US 09/890,456

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<150> IL 132846

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tggtaaaaga	tgaggaaact	gaggctcaga	gagggtgaagt	acctggccca	aggccacaca	1560
gccagaatct	tccacttgac	tcagatcaag	aaagtcagga	agcaagactt	ccagaaagag	1620
gcacagcact	tccgactgct	cgttgcccc	cacgaaggtc	actggaacgt	cttccctagc	1680
cagaccctgg	agctgaagg	cacggccagt	ccagacaaag	tgaccaagac	ataacaaaga	1740
cctaacagtt	gcagatatga	gctgtataat	tgttgttatt	atatattaat	aaataagaag	1800
ttgcataacc	atcaaaa					1817

<210> 9
 <211> 1622
 <212> DNA
 <213> Homo sapiens

<400> 9

ggacaaataa	aaaggaaaca	agcatgattg	tgagggcaga	ggagcgtggg	actgagtcag	60
gagactggtg	ctgtcatcgc	tgccctggtga	ctgacttgct	gtgtggccct	caggtgtaac	120
ttacctcttc	tgggcctcat	ttgtctaata	ataataatta	acgctgatac	catgatataa	180
atctgtacag	catttcactg	cttgattccc	taactgccct	gtgagataag	cgttaaggct	240
cagagacagt	ggcatgccc	gtgttgacac	gtaagtgtgt	ggtaaagccg	agattcaaac	300
tcagaccttc	tggccccctt	cctaggagag	catgcccagt	tgtctagcag	attctctttt	360

gcctgagtgg	cccagatgac	atctctttta	gagctagaaa	gaaggagaaa	tgagacaggg	420
tctttgggct	ggagcctcct	gggactaaca	tggcactggg	cggtttgcca	ggcccagaca	480
tgttctgcct	tttccatggg	aagagatact	cccccggcga	gagctggcac	ccctacttgg	540
agccacaagg	cctgatgtac	tgcttgcgct	gtacctgctc	agagggcgcc	catgtgagtt	600
gttaccgcct	ccactgtccg	cctgtccact	gccccagcc	tgtgacggag	ccacagcaat	660
gctgtcccaa	gtgtgtggaa	cctcacactc	cctctggact	ccgggccccca	ccaaagtcc	720
gccagcacia	cgggaccatg	taccaacacg	gagagatctt	cagtgcccat	gagctgttcc	780
cctcccgct	gcccaccag	tgtgtcctct	gcagctgcac	agagggccag	atctactgcg	840
gcctcacaac	ctgccccgaa	ccaggctgcc	cagcaccct	cccgtgcc	gactcctgct	900
gccaaagcctg	caaagatgag	gcaagtgage	aatcggatga	agaggacagt	gtgcagtcgc	960
tccatgggg	gagacatcct	caggatccat	gttccagtga	tgctgggaga	aagagaggcc	1020
cgggcacccc	agccccact	ggcctcagcg	ccctctgag	cttcatccct	cgccacttca	1080
gacccaagg	agcaggcagc	acaactgtca	agatcgtcct	gaaggagaaa	cataagaaa	1140
aggacaaagc	agaccctggc	cacagtgaga	tcagttctac	caggtgtccc	aaggcacccg	1200
gcccggctct	cgtccacaca	tcggtatccc	caagcccaga	caacctgcgt	cgctttgccc	1260
tggaacacga	ggcctcggac	ttggtggaga	tctacctctg	gaagctggta	aaagatgagg	1320
aaactgaggc	tcagagaggt	gaagtacctg	gcccaggcc	acacagccag	aatcttccac	1380
ttgactcaga	tcaagaaagt	caggaagcaa	gacttccaga	aagaggcaca	gcacttccga	1440
ctgctcgctg	gccccacga	aggtcactgg	aacgtcttcc	tagcccagac	cctggagctg	1500
aaggtcacgg	ccagtcaga	caaagtgacc	aagacataac	aaagacctaa	cagttgcaga	1560
tatgagctgt	ataattgttg	ttattatata	ttaataaata	agaagttgca	taaccatcaa	1620
aa						1622

<210> 10
 <211> 1567
 <212> DNA
 <213> Homo sapiens

<400> 10						
ggacaaataa	aaaggaaaca	agcatgattg	tgagggcaga	ggagcgtggg	actgagtcag	60
gagactgggtg	ctgtcatcgc	tgcttgggtga	ctgacttgc	gtgtggccct	caggtgtaac	120
ttacctctc	tgggcctcat	ttgtctaate	ataataatta	acgctgatac	catgatataa	180
atctgtacag	catttccactg	cttgattccc	taactgcct	gtgagataag	cgttaaggct	240
cagagacagt	ggcatgccc	gtgttgacac	gtaagtgtgt	ggtaaagccg	agattcaaac	300
tcagaccttc	tggccccctg	cctaggagag	catgcccagt	tgtctagcag	attctctttt	360
gcctgagtgg	cccagatgac	atctctttta	gagctagaaa	gaaggagaaa	tgagacaggg	420
tctttgggct	ggagcctcct	gggactaaca	tggcactggg	cggtttgcca	ggcccagaca	480
tgttctgcct	tttccatggg	aagagatact	cccccggcga	gagctggcac	ccctacttgg	540
agccacaagg	cctgatgtac	tgcttgcgct	gtacctgctc	agagggcgcc	catgtgagtt	600
gttaccgcct	ccactgtccg	cctgtccact	gccccagcc	tgtgacggag	ccacagcaat	660
gctgtcccaa	gtgtgtggaa	cctcacactc	cctctggact	ccgggccccca	ccaaagtcc	720
gccagcacia	cgggaccatg	taccaacacg	gagagatctt	cagtgcccat	gagctgttcc	780
cctcccgct	gcccaccag	tgtgtcctct	gcagctgcac	agagggccag	atctactgcg	840
gcctcacaac	ctgccccgaa	ccaggctgcc	cagcaccct	cccgtgcc	gactcctgct	900
gccaaagcctg	caaagatgag	gcaagtgage	aatcggatga	agaggacagt	gtgcagtcgc	960
tccatgggg	gagacatcct	caggatccat	gttccagtga	tgctgggaga	aagagaggcc	1020
cgggcacccc	agccccact	ggcctcagcg	ccctctgag	cttcatccct	cgccacttca	1080
gacccaagg	agcaggcagc	acaactgtca	agatcgtcct	gaaggagaaa	cataagaaa	1140
aggacaaagc	agaccctggc	cacagtgaga	tcagttctac	caggtgtccc	aaggcacccg	1200
gcccggctct	cgtccacaca	tcggtatccc	caagcccaga	caacctgcgt	cgctttgccc	1260
tggaacacga	ggcctcggac	ttggtggaga	tctacctctg	gaagctggta	aaaggaatct	1320
tccacttgac	tcagatcaag	aaagtcaagg	agcaagactt	ccagaaagag	gcacagcact	1380
tccgactgct	cgttgcccc	cacgaaggct	actggaacgt	cttcctagcc	cagacctgg	1440
agctgaaggt	cacggccagt	ccagacaaag	tgaccaagac	ataacaaaga	cctaacagtt	1500
gcagatatga	gctgtataat	tgttgttatt	atatattaat	aaataagaag	ttgcataacc	1560
atcaaaa						1567

<210> 11
 <211> 1202
 <212> DNA
 <213> Mouse

<220>
 <221> UNSURE
 <222> (1)..(1202)
 <223> n = a,c,g,t any unknown or other

<400> 11
 atttctctat tcttgatccc aactgtctct gcctaccac accagcccca aggtctnaga 60
 aagccctgga ggctggcttg ccaaatacctt gtcagtgtnt ttattgatta gtctgagaat 120
 atottagacc tcacccacaa ggttctgtgt ggagcctgtg ctctctgtct gtctgtctgt 180
 ctgtctgtct gtctgtctgt ctgcctgcct ctctctgtct gtctccgtct gtctctgtct 240
 ctctgtctgt ctctgtctgt ctctttctct ctgtctctct ctgtgtctct gtctctgtct 300
 ctgtctctct ctctctctca gaagtcctct agccttctct agcaggcgtc tcatgcagcc 360
 tgggttgggtg tcccagctgt ggcctatccc acagacagct ccacatcctg cttgtgttcc 420
 gcagagacat tcccaggatc catgctcgga gaggagaggc cccagcacgc cagcccccac 480
 cagcctcagc tcccctctgg gcttcatccn tcgccacttc cagtcagtag gaatgggcag 540
 cacaaccatc aagattatct tgaaggagaa acataaaaaa gcttgcacac acaatgggaa 600
 gacatactcc catggggagg tgtggcacc cactgtgtct tcctttggcc ccatgccctg 660
 catcctgtgc acatgtattg atggctacca ggactgccac cgtgtgacct gccccaccca 720
 atatccctgc agtcaaccca agaaagtggc tgggaagtgc tgcaagatct gccagagga 780
 cgaggcggaa gatgaccaca gtgaggtcat ttccaccggg tgtcccaagg taccaggcca 840
 gttccagggtg tacacgttgg catctccaag cccagacagc ctacaccgct ttgtcctgga 900
 gcatgaagcc tctgaccagg tagagatgta catttggaag ctgggtgaaag gaatttacca 960
 cttgggttcag atcaagagag tcaggaagca agatttccag aaagagggtc agaacttccg 1020
 gctgctcacc ggcacccatg aaggttactg gacogttttc ctagcccaga ttccagagct 1080
 gaaagttaca gccagcccag acaaagtgc caagacatta tagcaaggac ctaaagagtt 1140
 gcagatacga gttttatttg ttttgttatt atatattaat aaagaagtcg cattaccctt 1200
 tc 1202

<210> 12
 <211> 398
 <212> PRT
 <213> Homo sapiens

<220>
 <221> UNSURE
 <222> (1)..(398)
 <223> Xaa = any amino acid, unknown or other

<400> 12
 Arg Glu Pro Gln Gly Leu Met Tyr Cys Leu Arg Cys Thr Cys Ser Glu
 1 5 10 15
 Gly Ala His Val Ser Cys Tyr Arg Leu His Cys Pro Pro Val His Cys
 20 25 30
 Pro Gln Pro Val Thr Glu Pro Gln Gln Cys Cys Pro Lys Cys Val Glu
 35 40 45
 Pro His Thr Pro Ser Gly Leu Arg Ala Pro Pro Lys Ser Cys Gln His

50	55	60
Asn Gly Thr Met Tyr Gln His Gly Glu Ile Phe Ser Ala His Glu Leu 65 70 75 80		
Phe Pro Ser Arg Leu Pro Asn Gln Cys Val Leu Cys Ser Cys Thr Glu 85 90 95		
Gly Gln Ile Tyr Cys Gly Leu Thr Thr Cys Pro Glu Pro Gly Cys Pro 100 105 110		
Ala Pro Leu Pro Leu Pro Asp Ser Cys Cys Gln Ala Cys Lys Asp Glu 115 120 125		
Ala Ser Glu Gln Ser Asp Glu Glu Asp Ser Val Gln Ser Leu His Gly 130 135 140		
Val Arg His Pro Gln Asp Pro Cys Ser Ser Asp Ala Gly Arg Lys Arg 145 150 155 160		
Gly Pro Gly Thr Pro Ala Pro Thr Gly Leu Ser Ala Pro Leu Ser Phe 165 170 175		
Ile Pro Arg His Phe Arg Pro Lys Gly Ala Gly Ser Thr Thr Val Lys 180 185 190		
Ile Val Leu Lys Glu Lys His Xaa Lys Ala Cys Val His Gly Gly Lys 195 200 205		
Thr Tyr Ser His Gly Glu Val Trp His Pro Ala Phe Arg Ala Phe Gly 210 215 220		
Pro Cys Pro Cys Ile Leu Cys Thr Cys Glu Asp Gly Arg Gln Asp Cys 225 230 235 240		
Gln Arg Val Thr Cys Pro Thr Lys Tyr Pro Cys Arg His Pro Glu Lys 245 250 255		
Val Ala Gly Lys Cys Cys Lys Ile Cys Pro Glu Asp Lys Ala Asp Pro 260 265 270		
Gly His Ser Glu Ile Ser Ser Thr Arg Cys Pro Lys Ala Pro Gly Arg 275 280 285		
Val Leu Val His Thr Ser Val Ser Pro Ser Pro Asp Asn Leu Arg Arg 290 295 300		
Phe Ala Leu Glu His Glu Ala Ser Asp Leu Val Glu Ile Tyr Leu Trp 305 310 315 320		
Lys Leu Val Lys Asp Glu Glu Thr Glu Ala Gln Arg Gly Glu Val Pro 325 330 335		
Gly Pro Arg Pro His Ser Gln Asn Phe His Leu Thr Gln Ile Lys Lys 340 345 350		
Val Arg Lys Gln Asp Phe Gln Lys Glu Ala Gln His Phe Arg Leu Leu		

355 360 365
Ala Gly Pro His Glu Gly His Trp Asn Val Phe Leu Ala Gln Thr Leu
370 375 380

Glu Leu Lys Val Thr Ala Ser Pro Asp Lys Val Thr Lys Thr
385 390 395

<210> 13
<211> 539
<212> PRT
<213> Homo sapiens

<220>
<221> UNSURE
<222> (1)..(539)
<223> Xaa = any amino acid, unknown or other

<400> 13
Ser Pro Leu Pro Ser Ala Gly Pro Ser Phe Val Ser Pro Ser Leu Pro
1 5 10 15

Pro Phe Pro Ala Phe Ser Phe His Leu Ser Leu Leu Pro Thr Leu Asp
20 25 30

Leu Pro Ser Cys Pro Pro Phe Leu Pro Thr Ala Ala Ser Trp Pro Phe
35 40 45

Ser Asp Pro Ala Leu Ala Ala Asp Leu Leu Gly Ser Cys Gly Leu Ile
50 55 60

Cys Gly Pro Cys Xaa Ser Val Ser Phe Ser Ser Pro Val Leu Pro Thr
65 70 75 80

Pro Leu Pro Asp Gln Arg Pro Asp Pro Gly Glu Arg Met Val Pro Glu
85 90 95

Val Arg Val Leu Ser Ser Leu Leu Gly Leu Ala Leu Leu Trp Phe Pro
100 105 110

Leu Asp Ser His Ala Arg Ala Arg Pro Asp Met Phe Cys Leu Phe His
115 120 125

Gly Lys Arg Tyr Ser Pro Gly Glu Ser Trp His Pro Tyr Leu Glu Pro
130 135 140

Gln Gly Leu Met Tyr Cys Leu Arg Cys Thr Cys Ser Glu Gly Ala His
145 150 155 160

Val Ser Cys Tyr Arg Leu His Cys Pro Pro Val His Cys Pro Gln Pro
165 170 175

Val Thr Glu Pro Gln Gln Cys Cys Pro Lys Cys Val Glu Pro His Thr
180 185 190

Pro Ser Gly Leu Arg Ala Pro Pro Lys Ser Cys Gln His Asn Gly Thr

195					200					205					
Met	Tyr	Gln	His	Gly	Glu	Ile	Phe	Ser	Ala	His	Glu	Leu	Phe	Pro	Ser
210					215					220					
Arg	Leu	Pro	Asn	Gln	Cys	Val	Leu	Cys	Ser	Cys	Thr	Glu	Gly	Gln	Ile
225					230					235					240
Tyr	Cys	Gly	Leu	Thr	Thr	Cys	Pro	Glu	Pro	Gly	Cys	Pro	Ala	Pro	Leu
				245					250					255	
Pro	Leu	Pro	Asp	Ser	Cys	Cys	Gln	Ala	Cys	Lys	Asp	Glu	Ala	Ser	Glu
			260					265					270		
Gln	Ser	Asp	Glu	Glu	Asp	Ser	Val	Gln	Ser	Leu	His	Gly	Val	Arg	His
		275					280					285			
Pro	Gln	Asp	Pro	Cys	Ser	Ser	Asp	Ala	Gly	Arg	Lys	Arg	Gly	Pro	Gly
	290					295					300				
Thr	Pro	Ala	Pro	Thr	Gly	Leu	Ser	Ala	Pro	Leu	Ser	Phe	Ile	Pro	Arg
305					310					315					320
His	Phe	Arg	Pro	Lys	Gly	Ala	Gly	Ser	Thr	Thr	Val	Lys	Ile	Val	Leu
				325					330					335	
Lys	Glu	Lys	His	Xaa	Lys	Ala	Cys	Val	His	Gly	Gly	Lys	Thr	Tyr	Ser
			340					345					350		
His	Gly	Glu	Val	Trp	His	Pro	Ala	Phe	Arg	Ala	Phe	Gly	Pro	Cys	Pro
		355					360					365			
Cys	Ile	Leu	Cys	Thr	Cys	Glu	Asp	Gly	Arg	Gln	Asp	Cys	Gln	Arg	Val
	370					375					380				
Thr	Cys	Pro	Thr	Lys	Tyr	Pro	Cys	Arg	His	Pro	Glu	Lys	Val	Ala	Gly
385					390					395					400
Lys	Cys	Cys	Lys	Ile	Cys	Pro	Glu	Asp	Lys	Ala	Asp	Pro	Gly	His	Ser
				405					410					415	
Glu	Ile	Ser	Ser	Thr	Arg	Cys	Pro	Lys	Ala	Pro	Gly	Arg	Val	Leu	Val
			420					425					430		
His	Thr	Ser	Val	Ser	Pro	Ser	Pro	Asp	Asn	Leu	Arg	Arg	Phe	Ala	Leu
		435					440					445			
Glu	His	Glu	Ala	Ser	Asp	Leu	Val	Glu	Ile	Tyr	Leu	Trp	Lys	Leu	Val
	450					455					460				
Lys	Asp	Glu	Glu	Thr	Glu	Ala	Gln	Arg	Gly	Glu	Val	Pro	Gly	Pro	Arg
465					470					475					480
Pro	His	Ser	Gln	Asn	Phe	His	Leu	Thr	Gln	Ile	Lys	Lys	Val	Arg	Lys
				485					490					495	
Gln	Asp	Phe	Gln	Lys	Glu	Ala	Gln	His	Phe	Arg	Leu	Leu	Ala	Gly	Pro

500	505	510
His Glu Gly His Trp Asn Val Phe Leu Ala Gln Thr Leu Glu Leu Lys		
515	520	525
Val Thr Ala Ser Pro Asp Lys Val Thr Lys Thr		
530	535	

<210> 14
 <211> 388
 <212> PRT
 <213> Homo sapiens

<220>
 <221> UNSURE
 <222> (1)..(388)
 <223> Xaa = any amino acid, unknown or other

<400> 14
 Ile Ser Ser Trp Gly Gln Met Gln Asn His Gln Lys Ser Gly Leu Val
 1 5 10 15
 Asn Phe Ser Lys Asp Ser His Glu Thr Ser Phe Ser Ser Ser Ser Cys
 20 25 30
 Pro Ser Pro Thr Val Glu Pro His Thr Pro Ser Gly Leu Arg Ala Pro
 35 40 45
 Pro Lys Ser Cys Gln His Asn Gly Thr Met Tyr Gln His Gly Glu Ile
 50 55 60
 Phe Ser Ala His Glu Leu Phe Pro Ser Arg Leu Pro Asn Gln Cys Val
 65 70 75 80
 Leu Cys Ser Cys Thr Glu Gly Gln Ile Tyr Cys Gly Leu Thr Thr Cys
 85 90 95
 Pro Glu Pro Gly Cys Pro Ala Pro Leu Pro Leu Pro Asp Ser Cys Cys
 100 105 110
 Gln Ala Cys Lys Asp Glu Ala Ser Glu Gln Ser Asp Glu Glu Asp Ser
 115 120 125
 Val Gln Ser Leu His Gly Val Arg His Pro Gln Asp Pro Cys Ser Ser
 130 135 140
 Asp Ala Gly Arg Lys Arg Gly Pro Gly Thr Pro Ala Pro Thr Gly Leu
 145 150 155 160
 Ser Ala Pro Leu Ser Phe Ile Pro Arg His Phe Arg Pro Lys Gly Ala
 165 170 175
 Gly Ser Thr Thr Val Lys Ile Val Leu Lys Glu Lys His Xaa Lys Ala
 180 185 190
 Cys Val His Gly Gly Lys Thr Tyr Ser His Gly Glu Val Trp His Pro

195					200					205					
Ala	Phe	Arg	Ala	Phe	Gly	Pro	Cys	Pro	Cys	Ile	Leu	Cys	Thr	Cys	Glu
210						215					220				
Asp	Gly	Arg	Gln	Asp	Cys	Gln	Arg	Val	Thr	Cys	Pro	Thr	Lys	Tyr	Pro
225					230					235					240
Cys	Arg	His	Pro	Glu	Lys	Val	Ala	Gly	Lys	Cys	Cys	Lys	Ile	Cys	Pro
				245					250					255	
Glu	Asp	Lys	Ala	Asp	Pro	Gly	His	Ser	Glu	Ile	Ser	Ser	Thr	Arg	Cys
			260					265					270		
Pro	Lys	Ala	Pro	Gly	Arg	Val	Leu	Val	His	Thr	Ser	Val	Ser	Pro	Ser
		275					280					285			
Pro	Asp	Asn	Leu	Arg	Arg	Phe	Ala	Leu	Glu	His	Glu	Ala	Ser	Asp	Leu
	290					295					300				
Val	Glu	Ile	Tyr	Leu	Trp	Lys	Leu	Val	Lys	Asp	Glu	Glu	Thr	Glu	Ala
305					310					315					320
Gln	Arg	Gly	Glu	Val	Pro	Gly	Pro	Arg	Pro	His	Ser	Gln	Asn	Phe	His
				325					330					335	
Leu	Thr	Gln	Ile	Lys	Lys	Val	Arg	Lys	Gln	Asp	Phe	Gln	Lys	Glu	Ala
			340					345					350		
Gln	His	Phe	Arg	Leu	Leu	Ala	Gly	Pro	His	Glu	Gly	His	Trp	Asn	Val
		355					360					365			
Phe	Leu	Ala	Gln	Thr	Leu	Glu	Leu	Lys	Val	Thr	Ala	Ser	Pro	Asp	Lys
	370					375					380				
Val	Thr	Lys	Thr												
385															

<210> 15
 <211> 439
 <212> PRT
 <213> Homo sapiens

<220>
 <221> UNSURE
 <222> (1)..(439)
 <223> Xaa = any amino acid, unknown or other

<400> 15
 Asp Arg Val Phe Gly Leu Glu Pro Pro Gly Thr Asn Met Ala Leu Val
 1 5 10 15
 Gly Leu Pro Gly Pro Asp Met Phe Cys Leu Phe His Gly Lys Arg Tyr
 20 25 30
 Ser Pro Gly Glu Ser Trp His Pro Tyr Leu Glu Pro Gln Gly Leu Met

35					40					45					
Tyr	Cys	Leu	Arg	Cys	Thr	Cys	Ser	Glu	Gly	Ala	His	Val	Ser	Cys	Tyr
	50					55					60				
Arg	Leu	His	Cys	Pro	Pro	Val	His	Cys	Pro	Gln	Pro	Val	Thr	Glu	Pro
	65					70					75				80
Gln	Gln	Cys	Cys	Pro	Lys	Cys	Val	Glu	Pro	His	Thr	Pro	Ser	Gly	Leu
				85					90					95	
Arg	Ala	Pro	Pro	Lys	Ser	Cys	Gln	His	Asn	Gly	Thr	Met	Tyr	Gln	His
				100					105					110	
Gly	Glu	Ile	Phe	Ser	Ala	His	Glu	Leu	Phe	Pro	Ser	Arg	Leu	Pro	Asn
		115						120					125		
Gln	Cys	Val	Leu	Cys	Ser	Cys	Thr	Glu	Gly	Gln	Ile	Tyr	Cys	Gly	Leu
	130					135					140				
Thr	Thr	Cys	Pro	Glu	Pro	Gly	Cys	Pro	Ala	Pro	Leu	Pro	Leu	Pro	Asp
	145					150					155				160
Ser	Cys	Cys	Gln	Ala	Cys	Lys	Asp	Glu	Ala	Ser	Glu	Gln	Ser	Asp	Glu
				165					170					175	
Glu	Asp	Ser	Val	Gln	Ser	Leu	His	Gly	Val	Arg	His	Pro	Gln	Asp	Pro
			180						185					190	
Cys	Ser	Ser	Asp	Ala	Gly	Arg	Lys	Arg	Gly	Pro	Gly	Thr	Pro	Ala	Pro
			195					200					205		
Thr	Gly	Leu	Ser	Ala	Pro	Leu	Ser	Phe	Ile	Pro	Arg	His	Phe	Arg	Pro
	210					215					220				
Lys	Gly	Ala	Gly	Ser	Thr	Thr	Val	Lys	Ile	Val	Leu	Lys	Glu	Lys	His
	225					230					235				240
Xaa	Lys	Ala	Cys	Val	His	Gly	Gly	Lys	Thr	Tyr	Ser	His	Gly	Glu	Val
				245					250					255	
Trp	His	Pro	Ala	Phe	Arg	Ala	Phe	Gly	Pro	Cys	Pro	Cys	Ile	Leu	Cys
			260						265					270	
Thr	Cys	Glu	Asp	Gly	Arg	Gln	Asp	Cys	Gln	Arg	Val	Thr	Cys	Pro	Thr
		275						280					285		
Lys	Tyr	Pro	Cys	Arg	His	Pro	Glu	Lys	Val	Ala	Gly	Lys	Cys	Cys	Lys
	290					295					300				
Ile	Cys	Pro	Glu	Asp	Lys	Ala	Asp	Pro	Gly	His	Ser	Glu	Ile	Ser	Ser
	305					310					315				320
Thr	Arg	Cys	Pro	Lys	Ala	Pro	Gly	Arg	Val	Leu	Val	His	Thr	Ser	Val
				325					330					335	
Ser	Pro	Ser	Pro	Asp	Asn	Leu	Arg	Arg	Phe	Ala	Leu	Glu	His	Glu	Ala

340 345 350
 Ser Asp Leu Val Glu Ile Tyr Leu Trp Lys Leu Val Lys Asp Glu Glu
 355 360 365
 Thr Glu Ala Gln Arg Gly Glu Val Pro Gly Pro Arg Pro His Ser Gln
 370 375 380
 Asn Phe His Leu Thr Gln Ile Lys Lys Val Arg Lys Gln Asp Phe Gln
 385 390 395 400
 Lys Glu Ala Gln His Phe Arg Leu Leu Ala Gly Pro His Glu Gly His
 405 410 415
 Trp Asn Val Phe Leu Ala Gln Thr Leu Glu Leu Lys Val Thr Ala Ser
 420 425 430
 Pro Asp Lys Val Thr Lys Thr
 435

<210> 16
 <211> 549
 <212> PRT
 <213> Homo sapiens

<400> 16
 Thr Phe Pro Leu Ser Leu Ile Ala Ser Pro Phe Cys Trp Thr Phe Leu
 1 5 10 15
 Arg Leu Ser Ile Ser Pro Ser Phe Pro Arg Val Leu Phe Pro Pro Phe
 20 25 30
 Ser Ser Ser His Leu Arg Pro Pro Phe Leu Pro Ser Phe Pro Ala His
 35 40 45
 Arg Cys Phe Leu Ala Leu Leu Arg Pro Arg Ser Ser Ser Arg Pro Pro
 50 55 60
 Gly Val Cys Gly Leu Ile Cys Gly Pro Cys Ala Ser Val Ser Phe Ser
 65 70 75 80
 Ser Pro Phe Leu Pro Thr Pro Leu Pro Asp Gln Arg Pro Asp Pro Gly
 85 90 95
 Glu Arg Met Val Pro Glu Val Arg Val Leu Ser Ser Leu Leu Gly Leu
 100 105 110
 Ala Leu Leu Trp Phe Pro Leu Asp Ser His Ala Arg Ala Arg Pro Asp
 115 120 125
 Met Phe Cys Leu Phe His Gly Lys Arg Tyr Ser Pro Gly Glu Ser Trp
 130 135 140
 His Pro Tyr Leu Glu Pro Gln Gly Leu Met Tyr Cys Leu Arg Cys Thr
 145 150 155 160

Cys Ser Glu Gly Ala His Val Ser Cys Tyr Arg Leu His Cys Pro Pro
 165 170 175
 Val His Cys Pro Gln Pro Val Thr Glu Pro Gln Gln Cys Cys Pro Lys
 180 185 190
 Cys Val Glu Pro His Thr Pro Ser Gly Leu Arg Ala Pro Pro Lys Ser
 195 200 205
 Cys Gln His Asn Gly Thr Met Tyr Gln His Gly Glu Ile Phe Ser Ala
 210 215 220
 His Glu Leu Phe Pro Ser Arg Leu Pro Asn Gln Cys Val Leu Cys Ser
 225 230 235 240
 Cys Thr Glu Gly Gln Ile Tyr Cys Gly Leu Thr Thr Cys Pro Glu Pro
 245 250 255
 Gly Cys Pro Ala Pro Leu Pro Leu Pro Asp Ser Cys Cys Gln Ala Cys
 260 265 270
 Lys Asp Glu Ala Ser Glu Gln Ser Asp Glu Glu Asp Arg Val Gln Ser
 275 280 285
 Leu His Gly Val Arg His Pro Gln Asp Pro Cys Ser Ser Asp Ala Gly
 290 295 300
 Arg Lys Arg Gly Pro Gly Thr Pro Ala Pro Thr Gly Leu Ser Ala Pro
 305 310 315 320
 Leu Ser Phe Ile Pro Arg His Phe Ile Pro Lys Gly Ala Gly Ser Thr
 325 330 335
 Thr Val Lys Ile Val Leu Lys Glu Lys His Lys Lys Ala Cys Val His
 340 345 350
 Gly Gly Lys Thr Tyr Ser His Gly Glu Val Trp His Pro Ala Phe Arg
 355 360 365
 Ala Phe Gly Pro Leu Pro Cys Ile Leu Cys Thr Cys Glu Asp Gly Arg
 370 375 380
 Gln Asp Cys Gln Arg Val Thr Cys Pro Thr Glu Tyr Pro Cys Arg His
 385 390 395 400
 Pro Glu Lys Val Ala Gly Lys Cys Cys Lys Ile Cys Pro Glu Asp Lys
 405 410 415
 Ala Asp Pro Gly His Ser Glu Ile Ser Ser Thr Arg Cys Pro Lys Ala
 420 425 430
 Pro Gly Arg Val Leu Val His Thr Ser Val Ser Pro Ser Pro Asp Asn
 435 440 445
 Leu Arg Arg Phe Ala Leu Glu His Glu Ala Ser Asp Leu Val Glu Ile
 450 455 460

Tyr Leu Trp Lys Leu Val Lys Asp Glu Glu Thr Glu Ala Gln Arg Gly
465 470 475 480

Glu Val Pro Gly Pro Arg Pro His Ser Gln Asn Leu Pro Leu Asp Ser
485 490 495

Asp Gln Glu Ser Gln Glu Ala Arg Leu Pro Glu Arg Gly Thr Ala Leu
500 505 510

Pro Thr Ala Arg Trp Pro Pro Arg Arg Ser Leu Glu Arg Leu Pro Ser
515 520 525

Pro Asp Pro Gly Ala Glu Gly His Gly Gln Ser Arg Gln Ser Asp Gln
530 535 540

Asp Ile Thr Lys Thr
545

<210> 17

<211> 549

<212> PRT

<213> Homo sapiens

<400> 17

Thr Phe Pro Leu Ser Leu Ile Ala Ser Pro Phe Cys Trp Thr Phe Leu
1 5 10 15

Arg Leu Ser Ile Ser Pro Ser Phe Pro Arg Val Leu Phe Pro Pro Phe
20 25 30

Ser Ser Ser His Leu Arg Pro Pro Phe Leu Pro Ser Phe Pro Ala His
35 40 45

Arg Cys Phe Leu Ala Leu Leu Arg Pro Arg Ser Ser Ser Arg Pro Pro
50 55 60

Gly Val Cys Gly Leu Ile Cys Gly Pro Cys Ala Ser Val Ser Phe Ser
65 70 75 80

Ser Pro Phe Leu Pro Thr Pro Leu Pro Asp Gln Arg Pro Asp Pro Gly
85 90 95

Glu Arg Met Val Pro Glu Val Arg Val Leu Ser Ser Leu Leu Gly Leu
100 105 110

Ala Leu Leu Trp Phe Pro Leu Asp Ser His Ala Arg Ala Arg Pro Asp
115 120 125

Met Phe Cys Leu Phe His Gly Lys Arg Tyr Ser Pro Gly Glu Ser Trp
130 135 140

His Pro Tyr Leu Glu Pro Gln Gly Leu Met Tyr Cys Leu Arg Cys Thr
145 150 155 160

Cys Ser Glu Gly Ala His Val Ser Cys Tyr Arg Leu His Cys Pro Pro
165 170 175

Val His Cys Pro Gln Pro Val Thr Glu Pro Gln Gln Cys Cys Pro Lys
180 185 190
Cys Val Glu Pro His Thr Pro Ser Gly Leu Arg Ala Pro Pro Lys Ser
195 200 205
Cys Gln His Asn Gly Thr Met Tyr Gln His Gly Glu Ile Phe Ser Ala
210 215 220
His Glu Leu Phe Pro Ser Arg Leu Pro Asn Gln Cys Val Leu Cys Ser
225 230 235 240
Cys Thr Glu Gly Gln Ile Tyr Cys Gly Leu Thr Thr Cys Pro Glu Pro
245 250 255
Gly Cys Pro Ala Pro Leu Pro Leu Pro Asp Ser Cys Cys Gln Ala Cys
260 265 270
Lys Gly Glu Ala Ser Glu Gln Ser Asp Glu Glu Asp Ser Val Gln Ser
275 280 285
Leu His Gly Val Arg His Pro Gln Asp Pro Cys Ser Ser Asp Ala Gly
290 295 300
Arg Lys Arg Gly Pro Gly Thr Pro Ala Pro Thr Gly Leu Ser Ala Pro
305 310 315 320
Leu Ser Phe Ile Pro Arg His Phe Arg Pro Lys Gly Ala Gly Ser Thr
325 330 335
Thr Val Lys Ile Val Leu Lys Glu Lys His Lys Lys Ala Cys Val His
340 345 350
Gly Gly Lys Thr Tyr Ser His Gly Glu Val Trp His Pro Ala Phe Arg
355 360 365
Ala Phe Gly Pro Leu Pro Cys Ile Leu Cys Thr Cys Glu Asp Gly Arg
370 375 380
Gln Asp Cys Gln Arg Val Thr Cys Pro Thr Glu Tyr Pro Cys Arg His
385 390 395 400
Pro Glu Lys Val Ala Gly Lys Cys Cys Lys Ile Cys Pro Glu Asp Lys
405 410 415
Ala Asp Pro Gly His Ser Glu Ile Ser Ser Thr Arg Cys Pro Lys Ala
420 425 430
Pro Gly Arg Val Leu Val His Thr Ser Val Ser Pro Ser Pro Asp Asn
435 440 445
Leu Arg Arg Phe Ala Leu Glu His Glu Ala Ser Asp Leu Val Glu Ile
450 455 460
Tyr Leu Trp Lys Leu Val Lys Asp Glu Glu Thr Glu Ala Gln Arg Gly
465 470 475 480

Glu Val Pro Gly Pro Arg Pro His Ser Gln Asn Leu Pro Leu Asp Ser
485 490 495

Asp Gln Glu Ser Gln Glu Ala Arg Leu Pro Glu Arg Gly Thr Ala Leu
500 505 510

Pro Thr Ala Arg Trp Pro Pro Arg Arg Ser Leu Glu Arg Leu Pro Ser
515 520 525

Pro Asp Pro Gly Ala Glu Gly His Gly Gln Ser Arg Gln Ser Asp Gln
530 535 540

Asp Ile Thr Lys Thr
545

<210> 18

<211> 392

<212> PRT

<213> Homo sapiens

<400> 18

Ile Ser Ser Trp Gly Gln Met Gln Asn His Gln Lys Ser Gly Leu Val
1 5 10 15

Asn Phe Ser Lys Asp Ser His Glu Thr Ser Phe Ser Ser Ser Ser Cys
20 25 30

Pro Ser Pro Thr Ala Glu Pro His Thr Pro Ser Gly Leu Arg Ala Pro
35 40 45

Pro Lys Ser Cys Gln His Asn Gly Thr Met Tyr Gln His Gly Glu Ile
50 55 60

Phe Ser Ala His Glu Leu Phe Pro Ser Arg Leu Pro Asn Gln Cys Val
65 70 75 80

Leu Cys Ser Cys Thr Glu Gly Gln Ile Tyr Cys Gly Leu Thr Thr Cys
85 90 95

Pro Glu Pro Gly Cys Pro Ala Pro Leu Pro Leu Pro Asp Ser Cys Cys
100 105 110

Gln Ala Cys Lys Asp Glu Ala Ser Glu Gln Ser Asp Glu Glu Asp Ser
115 120 125

Val Gln Ser Leu His Gly Val Arg His Pro Gln Asp Pro Cys Ser Ser
130 135 140

Asp Ala Gly Arg Lys Arg Gly Pro Gly Thr Pro Ala Pro Thr Gly Leu
145 150 155 160

Ser Ala Pro Leu Ser Phe Ile Pro Arg His Phe Arg Pro Lys Gly Ala
165 170 175

Gly Ser Thr Thr Val Lys Ile Val Leu Lys Glu Lys His Lys Lys Ala

180					185					190					
Cys	Val	His	Gly	Gly	Lys	Thr	Tyr	Ser	His	Gly	Glu	Val	Trp	His	Pro
		195					200					205			
Ala	Phe	Arg	Ala	Phe	Gly	Pro	Leu	Pro	Cys	Ile	Leu	Cys	Thr	Cys	Glu
	210					215					220				
Asp	Gly	Arg	Gln	Asp	Cys	Gln	Arg	Val	Thr	Cys	Pro	Thr	Glu	Tyr	Pro
225					230					235					240
Cys	Arg	His	Pro	Glu	Lys	Val	Ala	Gly	Lys	Cys	Cys	Lys	Ile	Cys	Pro
				245					250					255	
Glu	Asp	Lys	Ala	Asp	Pro	Gly	His	Ser	Glu	Ile	Ser	Ser	Thr	Arg	Cys
			260					265					270		
Pro	Lys	Ala	Pro	Gly	Arg	Val	Leu	Val	His	Thr	Ser	Val	Ser	Pro	Ser
	275						280					285			
Pro	Asp	Asn	Leu	Arg	Arg	Phe	Ala	Leu	Glu	His	Glu	Ala	Ser	Asp	Leu
	290					295					300				
Val	Glu	Ile	Tyr	Leu	Trp	Lys	Leu	Val	Lys	Asp	Glu	Glu	Thr	Glu	Ala
305					310					315					320
Gln	Arg	Gly	Glu	Val	Pro	Gly	Pro	Arg	Pro	His	Ser	Gln	Asn	Leu	Pro
				325					330					335	
Leu	Asp	Ser	Asp	Gln	Glu	Ser	Gln	Glu	Ala	Arg	Leu	Pro	Glu	Arg	Gly
			340					345					350		
Thr	Ala	Leu	Pro	Thr	Ala	Arg	Trp	Pro	Pro	Arg	Arg	Ser	Leu	Glu	Arg
	355						360					365			
Leu	Pro	Ser	Pro	Asp	Pro	Gly	Ala	Glu	Gly	His	Gly	Gln	Ser	Arg	Gln
	370					375					380				
Ser	Asp	Gln	Asp	Ile	Thr	Lys	Thr								
385					390										

<210> 19
 <211> 443
 <212> PRT
 <213> Homo sapiens

<400> 19
 Asp Arg Val Phe Gly Leu Glu Pro Pro Gly Thr Asn Met Ala Leu Val
 1 5 10 15
 Gly Leu Pro Gly Pro Asp Met Phe Cys Leu Phe His Gly Lys Arg Tyr
 20 25 30
 Ser Pro Gly Glu Ser Trp His Pro Tyr Leu Glu Pro Gln Gly Leu Met
 35 40 45

Tyr Cys Leu Arg Cys Thr Cys Ser Glu Gly Ala His Val Ser Cys Tyr
 50 55 60
 Arg Leu His Cys Pro Pro Val His Cys Pro Gln Pro Val Thr Glu Pro
 65 70 75 80
 Gln Gln Cys Cys Pro Lys Cys Val Glu Pro His Thr Pro Ser Gly Leu
 85 90 95
 Arg Ala Pro Pro Lys Ser Cys Gln His Asn Gly Thr Met Tyr Gln His
 100 105 110
 Gly Glu Ile Phe Ser Ala His Glu Leu Phe Pro Ser Arg Leu Pro Asn
 115 120 125
 Gln Cys Val Leu Cys Ser Cys Thr Glu Gly Gln Ile Tyr Cys Gly Leu
 130 135 140
 Thr Thr Cys Pro Glu Pro Gly Cys Pro Ala Pro Leu Pro Leu Pro Asp
 145 150 155 160
 Ser Cys Cys Gln Ala Cys Lys Asp Glu Ala Ser Glu Gln Ser Asp Glu
 165 170 175
 Glu Asp Ser Val Gln Ser Leu His Gly Val Arg His Pro Gln Asp Pro
 180 185 190
 Cys Ser Ser Asp Ala Gly Arg Lys Arg Gly Pro Gly Thr Pro Ala Pro
 195 200 205
 Thr Gly Leu Ser Ala Pro Leu Ser Phe Ile Pro Arg His Phe Arg Pro
 210 215 220
 Lys Gly Ala Gly Ser Thr Thr Val Lys Ile Val Leu Lys Glu Lys His
 225 230 235 240
 Lys Lys Ala Cys Val His Gly Gly Lys Thr Tyr Ser His Gly Glu Val
 245 250 255
 Trp His Pro Ala Phe Arg Ala Phe Gly Pro Leu Pro Cys Ile Leu Cys
 260 265 270
 Thr Cys Glu Asp Gly Arg Gln Asp Cys Gln Arg Val Thr Cys Pro Thr
 275 280 285
 Glu Tyr Pro Cys Arg His Pro Glu Lys Val Ala Gly Lys Cys Cys Lys
 290 295 300
 Ile Cys Pro Glu Asp Lys Ala Asp Pro Gly His Ser Glu Ile Ser Ser
 305 310 315 320
 Thr Arg Cys Pro Lys Ala Pro Gly Arg Val Leu Val His Thr Ser Val
 325 330 335
 Ser Pro Ser Pro Asp Asn Leu Arg Arg Phe Ala Leu Glu His Glu Ala
 340 345 350

Ser Asp Leu Val Glu Ile Tyr Leu Trp Lys Leu Val Lys Asp Glu Glu
 355 360 365
 Thr Glu Ala Gln Arg Gly Glu Val Pro Gly Pro Arg Pro His Ser Gln
 370 375 380
 Asn Leu Pro Leu Asp Ser Asp Gln Glu Ser Gln Glu Ala Arg Leu Pro
 385 390 395 400
 Glu Arg Gly Thr Ala Leu Pro Thr Ala Arg Trp Pro Pro Arg Arg Ser
 405 410 415
 Leu Glu Arg Leu Pro Ser Pro Asp Pro Gly Ala Glu Gly His Gly Gln
 420 425 430
 Ser Arg Gln Ser Asp Gln Asp Ile Thr Lys Thr
 435 440

<210> 20
 <211> 378
 <212> PRT
 <213> Homo sapiens

<400> 20
 Asp Arg Val Phe Gly Leu Glu Pro Pro Gly Thr Asn Met Ala Leu Val
 1 5 10 15
 Gly Leu Pro Gly Pro Asp Met Phe Cys Leu Phe His Gly Lys Arg Tyr
 20 25 30
 Ser Pro Gly Glu Ser Trp His Pro Tyr Leu Glu Pro Gln Gly Leu Met
 35 40 45
 Tyr Cys Leu Arg Cys Thr Cys Ser Glu Gly Ala His Val Ser Cys Tyr
 50 55 60
 Arg Leu His Cys Pro Pro Val His Cys Pro Gln Pro Val Thr Glu Pro
 65 70 75 80
 Gln Gln Cys Cys Pro Lys Cys Val Glu Pro His Thr Pro Ser Gly Leu
 85 90 95
 Arg Ala Pro Pro Lys Ser Cys Gln His Asn Gly Thr Met Tyr Gln His
 100 105 110
 Gly Glu Ile Phe Ser Ala His Glu Leu Phe Pro Ser Arg Leu Pro Asn
 115 120 125
 Gln Cys Val Leu Cys Ser Cys Thr Glu Gly Gln Ile Tyr Cys Gly Leu
 130 135 140
 Thr Thr Cys Pro Glu Pro Gly Cys Pro Ala Pro Leu Pro Leu Pro Asp
 145 150 155 160
 Ser Cys Cys Gln Ala Cys Lys Asp Glu Ala Ser Glu Gln Ser Asp Glu
 165 170 175

Glu Asp Ser Val Gln Ser Leu His Gly Val Arg His Pro Gln Asp Pro
 180 185 190
 Cys Ser Ser Asp Ala Gly Arg Lys Arg Gly Pro Gly Thr Pro Ala Pro
 195 200 205
 Thr Gly Leu Ser Ala Pro Leu Ser Phe Ile Pro Arg His Phe Arg Pro
 210 215 220
 Lys Gly Ala Gly Ser Thr Thr Val Lys Ile Val Leu Lys Glu Lys His
 225 230 235 240
 Lys Lys Glu Asp Lys Ala Asp Pro Gly His Ser Glu Ile Ser Ser Thr
 245 250 255
 Arg Cys Pro Lys Ala Pro Gly Arg Val Leu Val His Thr Ser Val Ser
 260 265 270
 Pro Ser Pro Asp Asn Leu Arg Arg Phe Ala Leu Glu His Glu Ala Ser
 275 280 285
 Asp Leu Val Glu Ile Tyr Leu Trp Lys Leu Val Lys Asp Glu Glu Thr
 290 295 300
 Glu Ala Gln Arg Gly Glu Val Pro Gly Pro Arg Pro His Ser Gln Asn
 305 310 315 320
 Leu Pro Leu Asp Ser Asp Gln Glu Ser Gln Glu Ala Arg Leu Pro Glu
 325 330 335
 Arg Gly Thr Ala Leu Pro Thr Ala Arg Trp Pro Pro Arg Arg Ser Leu
 340 345 350
 Glu Arg Leu Pro Ser Pro Asp Pro Gly Ala Glu Gly His Gly Gln Ser
 355 360 365
 Arg Gln Ser Asp Gln Asp Ile Thr Lys Thr
 370 375

<210> 21
 <211> 356
 <212> PRT
 <213> Homo sapiens

<400> 21
 Asp Arg Val Phe Gly Leu Glu Pro Pro Gly Thr Asn Met Ala Leu Val
 1 5 10 15
 Gly Leu Pro Gly Pro Asp Met Phe Cys Leu Phe His Gly Lys Arg Tyr
 20 25 30
 Ser Pro Gly Glu Ser Trp His Pro Tyr Leu Glu Pro Gln Gly Leu Met
 35 40 45
 Tyr Cys Leu Arg Cys Thr Cys Ser Glu Gly Ala His Val Ser Cys Tyr

50					55					60					
Arg	Leu	His	Cys	Pro	Pro	Val	His	Cys	Pro	Gln	Pro	Val	Thr	Glu	Pro
65					70					75					80
Gln	Gln	Cys	Cys	Pro	Lys	Cys	Val	Glu	Pro	His	Thr	Pro	Ser	Gly	Leu
				85					90					95	
Arg	Ala	Pro	Pro	Lys	Ser	Cys	Gln	His	Asn	Gly	Thr	Met	Tyr	Gln	His
			100					105					110		
Gly	Glu	Ile	Phe	Ser	Ala	His	Glu	Leu	Phe	Pro	Ser	Arg	Leu	Pro	Asn
		115					120					125			
Gln	Cys	Val	Leu	Cys	Ser	Cys	Thr	Glu	Gly	Gln	Ile	Tyr	Cys	Gly	Leu
	130					135					140				
Thr	Thr	Cys	Pro	Glu	Pro	Gly	Cys	Pro	Ala	Pro	Leu	Pro	Leu	Pro	Asp
145					150					155					160
Ser	Cys	Cys	Gln	Ala	Cys	Lys	Asp	Glu	Ala	Ser	Glu	Gln	Ser	Asp	Glu
			165						170					175	
Glu	Asp	Ser	Val	Gln	Ser	Leu	His	Gly	Val	Arg	His	Pro	Gln	Asp	Pro
			180					185					190		
Cys	Ser	Ser	Asp	Ala	Gly	Arg	Lys	Arg	Gly	Pro	Gly	Thr	Pro	Ala	Pro
			195				200					205			
Thr	Gly	Leu	Ser	Ala	Pro	Leu	Ser	Phe	Ile	Pro	Arg	His	Phe	Arg	Pro
	210					215					220				
Lys	Gly	Ala	Gly	Ser	Thr	Thr	Val	Lys	Ile	Val	Leu	Lys	Glu	Lys	His
225					230					235					240
Lys	Lys	Glu	Asp	Lys	Ala	Asp	Pro	Gly	His	Ser	Glu	Ile	Ser	Ser	Thr
			245						250					255	
Arg	Cys	Pro	Lys	Ala	Pro	Gly	Arg	Val	Leu	Val	His	Thr	Ser	Val	Ser
			260					265					270		
Pro	Ser	Pro	Asp	Asn	Leu	Arg	Arg	Phe	Ala	Leu	Glu	His	Glu	Ala	Ser
			275				280					285			
Asp	Leu	Val	Glu	Ile	Tyr	Leu	Trp	Lys	Leu	Val	Lys	Gly	Ile	Phe	His
	290					295					300				
Leu	Thr	Gln	Ile	Lys	Lys	Val	Arg	Lys	Gln	Asp	Phe	Gln	Lys	Glu	Ala
305					310					315					320
Gln	His	Phe	Arg	Leu	Leu	Ala	Gly	Pro	His	Glu	Gly	His	Trp	Asn	Val
			325						330					335	
Phe	Leu	Ala	Gln	Thr	Leu	Glu	Leu	Lys	Val	Thr	Ala	Ser	Pro	Asp	Lys
			340					345					350		
Val	Thr	Lys	Thr												

<210> 22
 <211> 397
 <212> PRT
 <213> Mouse

<220>
 <221> UNSURE
 <222> (1)..(397)
 <223> Xaa = any amino acid, unknown or other

<400> 22
 Phe Leu Tyr Ser Ser His Thr Ala Leu Pro Thr His Thr Ser Pro Lys
 1 5 10 15
 Val Xaa Glu Ser Pro Gly Gly Trp Leu Ala Lys Ser Leu Ser Val Xaa
 20 25 30
 Leu Leu Ile Ser Leu Arg Ile Ser Thr Ser Pro Thr Arg Phe Cys Val
 35 40 45
 Glu Pro Val Leu Ser Val Cys Leu Ser Val Cys Leu Ser Val Cys Leu
 50 55 60
 Ser Ala Cys Leu Ser Leu Ser Val Ser Val Cys Leu Cys Leu Ser Val
 65 70 75 80
 Cys Leu Cys Leu Ser Leu Ser Leu Cys Leu Ser Leu Cys Leu Cys Leu
 85 90 95
 Cys Leu Cys Leu Ser Leu Ser Leu Arg Ser Pro Leu Ala Phe Ser Ser
 100 105 110
 Arg Arg Leu Met Gln Pro Gly Trp Cys Ser Gln Leu Trp Pro Ile Pro
 115 120 125
 Gln Thr Ala Pro His Pro Ala Cys Cys Ser Gln Arg His Ser Gln Asp
 130 135 140
 Pro Cys Ser Glu Arg Arg Gly Pro Ser Thr Pro Ala Pro Thr Ser Leu
 145 150 155 160
 Ser Ser Pro Leu Gly Phe Ile Xaa Arg His Phe Gln Ser Val Gly Met
 165 170 175
 Gly Ser Thr Thr Ile Lys Ile Ile Leu Lys Glu Lys His Lys Lys Ala
 180 185 190
 Cys Thr His Asn Gly Lys Thr Tyr Ser His Gly Glu Val Trp His Pro
 195 200 205
 Thr Val Leu Ser Phe Gly Pro Met Pro Cys Ile Leu Cys Thr Cys Ile
 210 215 220
 Asp Gly Tyr Gln Asp Cys His Arg Val Thr Cys Pro Thr Gln Tyr Pro

225		230		235		240
Cys Ser Gln Pro Lys Lys Val Ala Gly Lys Cys Cys Lys Ile Cys Pro						
	245			250		255
Glu Asp Glu Ala Glu Asp Asp His Ser Glu Val Ile Ser Thr Arg Cys						
	260		265			270
Pro Lys Val Pro Gly Gln Phe Gln Val Tyr Thr Leu Ala Ser Pro Ser						
	275		280			285
Pro Asp Ser Leu His Arg Phe Val Leu Glu His Glu Ala Ser Asp Gln						
	290		295			300
Val Glu Met Tyr Ile Trp Lys Leu Val Lys Gly Ile Tyr His Leu Val						
305		310		315		320
Gln Ile Lys Arg Val Arg Lys Gln Asp Phe Gln Lys Glu Val Gln Asn						
	325		330			335
Phe Arg Leu Leu Thr Gly Thr His Glu Gly Tyr Trp Thr Val Phe Leu						
	340		345			350
Ala Gln Ile Pro Glu Leu Lys Val Thr Ala Ser Pro Asp Lys Val Thr						
	355		360			365
Lys Thr Leu Gln Gly Pro Lys Glu Leu Gln Ile Arg Val Leu Leu Val						
	370		375			380
Leu Leu Leu Tyr Ile Asn Lys Glu Val Ala Leu Pro Phe						
385		390		395		

<210> 23
 <211> 21
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> DNA sense primer

<400> 23
 gaaagcctgt gtgcatggcg g

21

<210> 24
 <211> 23
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> DNA anti-sense primer

<400> 24
 agctcatatc tgcaactgtt agg

23